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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/642,405

DATE: 07/05/2001
TIME: 16:38:23

Input Set : A:\20413y.txt
Output Set: N:\CRF3\07032001\I642405.raw

4 <110> APPLICANT: Neeper, Michael P.
5 McClements, William L.
6 Jansen, Kathrin U.
7 Schultz, Loren D.
8 Chen, Ling
9 Wang, Xin-Min
11 <120> TITLE OF INVENTION: SYNTHETIC HUMAN PAPILLOMAVIRUS GENES
14 <130> FILE REFERENCE: 20413Y
16 <140> CURRENT APPLICATION NUMBER: 09/642,405
C--> 17 <141> CURRENT FILING DATE: 2001-06-21
19 <150> PRIOR APPLICATION NUMBER: PCT/US00/22932
20 <151> PRIOR FILING DATE: 2000-08-21
22 <150> PRIOR APPLICATION NUMBER: 60/210,143
23 <151> PRIOR FILING DATE: 2000-06-07
25 <150> PRIOR APPLICATION NUMBER: 60/150,728
26 <151> PRIOR FILING DATE: 1999-08-25
28 <160> NUMBER OF SEQ ID NOS: 150
30 <170> SOFTWARE: FastSEQ for Windows Version 4.0
32 <210> SEQ ID NO: 1
33 <211> LENGTH: 1518
34 <212> TYPE: DNA
35 <213> ORGANISM: Artificial Sequence
37 <220> FEATURE:
38 <223> OTHER INFORMATION: Codon-Optimized HPV16 L1
40 <400> SEQUENCE: 1

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43	cgcctgctgg	ccgtggccca	cccctacttc	cccatcaaga	agcccaacaa	caacaagatc	180
44	ctgggtccca	aggtgagcgg	cctgcagttac	cgcgtgttcc	gcatccaccc	gcccggacccc	240
45	aacaagttcg	gttcccccg	cacaagttc	tacaaccccg	acacccagcg	cctgggttgg	300
46	gcctgcgtgg	gcgtggagggt	ggggccggcgc	cagccctgg	gcgtgggcatt	cagccggccac	360
47	ccctgtgtca	acaagctgg	cgacaccgg	aacgcggcag	cctacgcggc	caacgcggc	420
48	gtggacaacc	gctggatgtcat	cagcatggac	tacaaggaga	cccagctgt	cctgatcgcc	480
49	tgcaaggctc	ccatcgccg	gcactggggc	aaggcagcc	cctgcaccaa	cgtggccgt	540
50	aaccccgccg	actgcctcc	cctggagctg	atcaacaccg	tgatccagga	cggcgacatg	600
51	gtggacaccg	gttccggcgc	catggacttc	accaccctgc	aggccaacaa	gagcgaggt	660
52	ccctggaca	tctgcaccag	catctgtcaag	tacccgact	acatcaagat	gttggatcgag	720
53	ccctacggcg	acagcctgtt	cttctacctg	cgccgcgagc	agatgttgcgt	gcgcacactg	780
54	ttcaaccggcg	ccggccgcgt	gggcgagaac	gtgcccgcgc	acctgtacat	caagggcagc	840
55	ggcagcaccg	ccaaacctggc	cagcagcaac	tacttccca	ctcccgccgg	cagcatggtg	900
56	accagcgacg	cccaaataatctt	caacaagccc	tactggctgc	agcgcgcccc	ggccacaac	960
57	aacgcacatct	gttggggca	ccagctgttc	gtgaccgtgg	tggacaccac	ccgcacgcacc	1020
58	aacatgagcc	tgtgcgcgc	catcagcacc	agcgagatca	cctacaagaa	caccaacttc	1080
59	aaggagtacc	tgcgccacgg	cgaggagatc	gacctgcagt	tcatcttcca	gctgtgcaag	1140
60	atcaccctga	cgccgcacgt	gatgacccatc	atccacagca	tgaacagcac	catccctggag	1200
61	gactggaact	tcggcctgca	gccccctccc	ggccgttaccc	tggaggacac	ctaccgcttc	1260
62	gtgaccagcc	aggccatcgc	ctggcagaag	cacaccctc	ccgctcccaa	ggaggatccc	1320

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63 ctgaagaagt acacccctcg ggaggtgaac ctgaaggaga agttcagcgc cgacccggac 1380
 64 cagttccccc tggccgc当地 gttcctgctg caggccggcc tgaaggccaa gcccaagttc 1440
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 69 <211> LENGTH: 1950
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 71 <213> ORGANISM: Artificial Sequence
 73 <220> FEATURE:
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 76 <400> SEQUENCE: 2
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 78 gtggaggccg tgggtggagaa gaagaccggc gacgccc当地 gcgacgacga gaacgagaac 120
 79 gacagc当地 cccggc当地 cctgggtggac ttc当地tgc当地 acgacaacga ctaccctgacc 180
 80 caggccgaga cccggc当地 cccggc当地 ccacccctg ttc当地ccggcc aggaggccaa gcagcaccgc 240
 81 gacgccc当地 aggtgctgaa ggc当地aactac ctggggc当地 ccctgagc当地 catcagcggc 300
 82 tgc当地tgc当地 acaacatc当地 cccccctg aaggccatct gcatc当地gagaa gcagagccgc 360
 83 gccc当地aaggc gccc当地tgg cggagc当地 gagc当地ggc当地 acggcaacac cgagggtggag 420
 84 accc当地c当地 gtc当地tgc当地ggt ggaggccgc caccggc当地 agacccctg cagcc当地gtac 480
 85 agccggc当地 gccggc当地ggg ctgc当地ggc当地 tacagc当地ggc当地 gcagcggc当地 cgaggggc当地 540
 86 agc当地gcc当地 acaccatctg ccagaccctt ctgaccaaca tcctg当地aaactg gctg当地agacc 600
 87 agcaacgcca aggccgccat gctggccaa ttc当地aggagc tgc当地ggc当地 gagttc当地g 660
 88 gagctggc当地 gccc当地tccaa gagcaacaag agcacctgct ggc当地actggc当地 catcggcc 720
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 90 tacctgc当地 cccaggccct ggc当地tgc当地 cccggc当地tgg tgggtgc当地 tgc当地ggc当地 840
 91 tacaagtgc当地 gcaagaaccg cgagaccatc gagaagctgc tgagcaagct gctgtgc当地gt 900
 92 agccccc当地atgt gcatgatgat cggccctcc aagcttgc当地 gcaccggccgc cggccctgta 960
 93 tggtaacaaga cccggc当地atc当地gca acacatc当地gagc gaggtgtacg ggc当地acccccc cgagtggatc 1020
 94 cagc当地gcc当地aga cccgtgctgca gcaacgcttcc aacgactgca ccttgc当地gact gaggccatg 1080
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 96 cagctggccg acaccaacag caacccggc当地 gccttcc当地tga agagcaacag ccaggccaaag 1200
 97 atcgtgaaagg actgc当地ccac catgtgc当地c当地 cactacaaggc ggc当地ccgagaa gaagc当地agatg 1260
 98 agcatgagcc agtggatcaa gtaccctgc gaccggc当地tgg acgacccggc当地 cgaccgcaag 1320
 99 cagatgtgca tggccctgc当地 ctaccaggc当地tggattca tgagcttccct gaccggccctg 1380
 100 aagc当地gttcc tgc当地aggccat ccccaagaag aactgc当地atcc tgc当地tgc当地gg cggcc当地caac 1440
 101 accgacaaga gccttgc当地tgg catgaccc当地tgc当地 atgaagttcc tgc当地aggccag cgtgatctgc 1500
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 104 gacggcaacc tggtaacgat ggacgtgaaag caccggccccc tggtaacgact gaaatggccct 1680
 105 cccctgctgca tcaccaggc当地 catcaacgccc ggc当地accggaca gccgctggcc ctaccctgc当地ac 1740
 106 aaccggccctg tggtaacgat cttccccc当地tgg gatcccttccct tc当地gacgagaa cggtaacc 1800
 107 gtgtacgagc tgaacgacaa gaaatggaaag agcttccctca gccgacccctg gaccggccctg 1860
 108 agcctgc当地acg aggacgagga caaggagaac gacggccgaca gcctgccc当地ac cttcaagtgca 1920
 109 gtgagcggcc agaacaccaa caccctgtaa 1950
 111 <210> SEQ ID NO: 3
 112 <211> LENGTH: 1098
 113 <212> TYPE: DNA
 114 <213> ORGANISM: Artificial Sequence
 116 <220> FEATURE:

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117 <223> OTHER INFORMATION: Mutant, Codon-Optimized HPV16 E2
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122 gccatctact acaaggcccg cgagatggc ttcaagcaca tcaaccacca ggtggtgccc 180
123 accctggccg tgagcaagaa caaggccctg caggccgcg agctgcacgt gaccctggag 240
124 accatctaca acagccagta cagcaacgag aagtggaccc tgcagacgt gagcctggag 300
125 gtgtacctga ccgccccccac cggtgcac aagaagcagc gctacaccgt ggaggtgcag 360
126 ttgcacggcg acatctgcaa caccatgcac tacaccaact ggaccacat ctacatctgc 420
127 gaggaggcca gcgtgaccgt ggtggaggc caggtggact actacggct gtactacgtg 480
128 cacgaggcca tccgcaccta cttcgtgcag ttcaaggacg acgcccggaa gtacagcaag 540
129 aacaagggtgt gggaggtgca cggccggggc caggtgatcc tggcccccac cagcgtgttc 600
130 agcagcaacg aggtgagcag ccccgagacc atccgcgc acctggccaa ccacagcgcc 660
131 gccaccacaca ccaaggccgt ggcctgggc accgaggaga cccagaccac catccagcgc 720
132 ccccgccagcg agcccgacac cggcaaccccc tgccacacca ccaagctgct gcaccgcgac 780
133 agcgtggaca gcgccccat cctgaccgc ttcaacagcga gcccacaaggg ccgcataac 840
134 tgcacacagca acaccacccc catcgtgcac ctgaagggcg acgccaacac cctgaagtgc 900
135 ctgcgcgtacc gcttcaagaa gcactgcaag ctgtacaccc cggtgagcag cacctggcac 960
136 tggaccggcc acaacgtgaa gcacaagagc gccatcgta ccctgaccta cgacagcgag 1020
137 tggcagcgcg accagttct gagccaggtg aagatccccca agaccatcac cgtgagcacc 1080
138 ggcttcatga gcatctaa 1098
140 <210> SEQ ID NO: 4
141 <211> LENGTH: 297
142 <212> TYPE: DNA
143 <213> ORGANISM: Artificial Sequence
145 <220> FEATURE:
146 <223> OTHER INFORMATION: Mutant, Codon-Optimized HPV16 E7
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150 gaccctgtacg gctacggcca gctgaacgc acgacgcagg aggaggacga gatcgacggc 120
151 cccgcccccc agggcgagcc cgaccgcgc cactacaaca tcgtgacctt ctgtgcaag 180
152 tgcacacagca ccctgcgcct gtgcgtgcag agcaccacac tggacatccg caccctggag 240
153 gacctgctga tggcacccct gggcatcgta tgcccatct gcagccagaa gccctaa 297
155 <210> SEQ ID NO: 5
156 <211> LENGTH: 297
157 <212> TYPE: DNA
158 <213> ORGANISM: Artificial Sequence
160 <220> FEATURE:
161 <223> OTHER INFORMATION: Codon-Optimized HPV6a E7
163 <400> SEQUENCE: 5
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165 gtggccctgc actgctacga gcagctggc gacagcagcg aggacgagggt ggacgaggtg 120
166 gacggccagg acagccagcc cctgaagcag cacttccaga tcgtgacctg ctgtgcggc 180
167 tgcacacagca acgtgcgcct ggtggcgcg tgacccgaga ccgacatccg cgaggtgcag 240
168 cagctcctgc tgggtacccct gaaacatcgta tgcccatct gcgtcccaa gacctaa 297
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171 <211> LENGTH: 318
172 <212> TYPE: DNA
173 <213> ORGANISM: Artificial Sequence

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175 <220> FEATURE:
 176 <223> OTHER INFORMATION: Codon-Optimized HPV18 E7
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 181 atcgacggcg tgaaccacca gcacctgcc gctgcgaggc ccgagcccca gcgccacacc 180
 182 atgctgtca tgcgtcgtcaa gtgcgaggcc cgcatcgagc tgggtgtgaa gagcagcgct 240
 183 gacgacctgc gcgccttcca gcagctgttc ctgaacaccc ttagcttcgt tgcccccgtgg 300
 184 tgcgccagcc agcagtaa 318
 186 <210> SEQ ID NO: 7
 187 <211> LENGTH: 1107
 188 <212> TYPE: DNA
 189 <213> ORGANISM: Artificial Sequence
 191 <220> FEATURE:
 192 <223> OTHER INFORMATION: Codon-Optimized HPV6a E2
 194 <400> SEQUENCE: 7
 195 atggaggcca tcgccaagcg cctggacgcc tgccaggagc agctgctgaa gctgtacgag 60
 196 gagaacagca ccgacctgca caagcacgtg ctgcactgaa agtgcacatgc ccacgagagc 120
 197 gtgtctgtgt acaaggccaa gcagatgggc ctgagccaca tcggcatgca ggtgggtgcct 180
 198 cctctgaagg tgagcgaggc caagggccac aacgcacatcg agatgcacatgc acacatcgag 240
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 200 atgtggcaga cccctcccaa gcgcgtgttc aagaagcgcg gcaagaccgt ggaggtgaag 360
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 204 accaaggact gggagggtgt ctacggcagc accgtgtatct gcagccccgc tagcgtgagc 600
 205 agecaccaccc aggagggtgag catccccgag agcaccaccc acactccccgc ccagaccgc 660
 206 accctggta gcagcagcac caaggaggac gccgtgcaga cccctccctcg caagcgcgc 720
 207 cggccgtgc agcagagccccc ctgcaacgcctt ctgtgcgtgg cccacatcg cccctggat 780
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 209 aacagcagcg ccactcccat cgtcagttc cagggcgaga gcaactgcct gaagtgccttc 900
 210 cgctaccgc tgaacgtatcg ccaccgcac ctgttcgacc tgatcagcag cacctggcacc 960
 211 tggggccagca gcaaggctcc ccacaaggac gccatgtga cctgtgaccta cgacagcgag 1020
 212 gacgacgcgc agcagttcct ggacgtgggtt aagatccctc ccaccatcg ccacaagctg 1080
 213 ggcttcatga gcctgcaccc gctgtaa 1107
 215 <210> SEQ ID NO: 8
 216 <211> LENGTH: 1098
 217 <212> TYPE: DNA
 218 <213> ORGANISM: Artificial Sequence
 220 <220> FEATURE:
 221 <223> OTHER INFORMATION: Codon-Optimized HPV18 E2
 223 <400> SEQUENCE: 8
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 225 gaccactacg agaacgcacag caaggacatc gacagccaga tccagtactg gcagctgtatc 120
 226 cgctgggaga acgccccatctt ctgcgtgtgtt cgcgagcaccg ggatcccgac cctgaaccac 180
 227 cagggtgtgc ccgccttacaa catcagcaag agcaggcccc acaaggccat cgacgtgcag 240
 228 atggccctgc agggccctggc ccagagcgcc tacaaggaccc aggactggac cctgcaggac 300
 229 acctgcgagg agctgtggaa caccgagcccc accccactgt tcaagaaggg aggcccagacc 360
 230 gtgcaggtgtt acttcgcacgg caacaaggac aactgcacatgc actacgtggc ctgggacagc 420

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231	gtgtactaca tgaccgacgc cggcacctgg gacaagaccg ccacctgcgt gagccaccgc	480
232	ggcctgtact acgtgaagga gggctacaac accttctaca tcgagttcaa gagcgagtgc	540
233	gagaagtacg gcaacaccgg cacctggag gtgcacttcg gcaacaacgt gatcgactgc	600
234	aacgacagca tgtcagcac cagcagcacc accgtgagcg ccacccagct ggtgaagcag	660
235	ctgcagcaca ctcccagccc ctacagcagc accgtgagcg tgggcaccgc caagacctac	720
236	gcccagacca gcgccgcccc tcgcctggc cactgcggcc tggccgagaa gcagcactgc	780
237	gggcccgtga accctctgt gggccccc accgcccaccc gcaacaacaa gcggcgaag	840
238	ctgtcagcg gcaacaccac tcccattatc caccgtgaaagg gcgaccgcaa cagcctgaag	900
239	tgcctgcgtt accgcctgcg caagcacagc gaccactacc gcgacatcag cagcacctgg	960
240	cactggaccg gcgccggaa cgagaagacc ggcattctga ccgtgaccta ccacagcgag	1020
241	acccagcgca ccaagtctt gaacaccgtg gccatcccc acagcgtgca gatcctggtg	1080
242	ggttacatga ccatgtaa	1098
244	<210> SEQ ID NO: 9	
245	<211> LENGTH: 129	
246	<212> TYPE: DNA	
247	<213> ORGANISM: Artificial Sequence	
249	<220> FEATURE:	
250	<223> OTHER INFORMATION: Codon-Optimized HPV16 L1 fragment	
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254	gttggtagca ccgacgagta cgtggccccc accaacatct actaccacgc cggcaccagc	120
255	cgccctgtgt	129
257	<210> SEQ ID NO: 10	
258	<211> LENGTH: 129	
259	<212> TYPE: DNA	
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263	<223> OTHER INFORMATION: Codon-Optimized HPV16 L1 fragment	
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267	gacaccccago gcttgggtgt ggcctgcgtt ggcgtggagg tggccgcgg ccagccccctg	120
268	ggcgtgggc	129
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271	<211> LENGTH: 129	
272	<212> TYPE: DNA	
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275	<220> FEATURE:	
276	<223> OTHER INFORMATION: Codon-Optimized HPV16 L1 fragment	
278	<400> SEQUENCE: 11	
279	gagtgcata gcatggacta caagcagacc cagctgtgcc tgatcggtgtt caagcctccc	60
280	atcggcgagc actggggcaaa gggcagcccc tgcaccaacg tggccgtt gcccggcgac	120
281	tgcctcccc	129
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284	<211> LENGTH: 132	
285	<212> TYPE: DNA	
286	<213> ORGANISM: Artificial Sequence	
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289	<223> OTHER INFORMATION: Codon-Optimized HPV16 L1 fragment	
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VERIFICATION SUMMARY
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L:17 M:271 C: Current Filing Date differs, Replaced Current Filing Date